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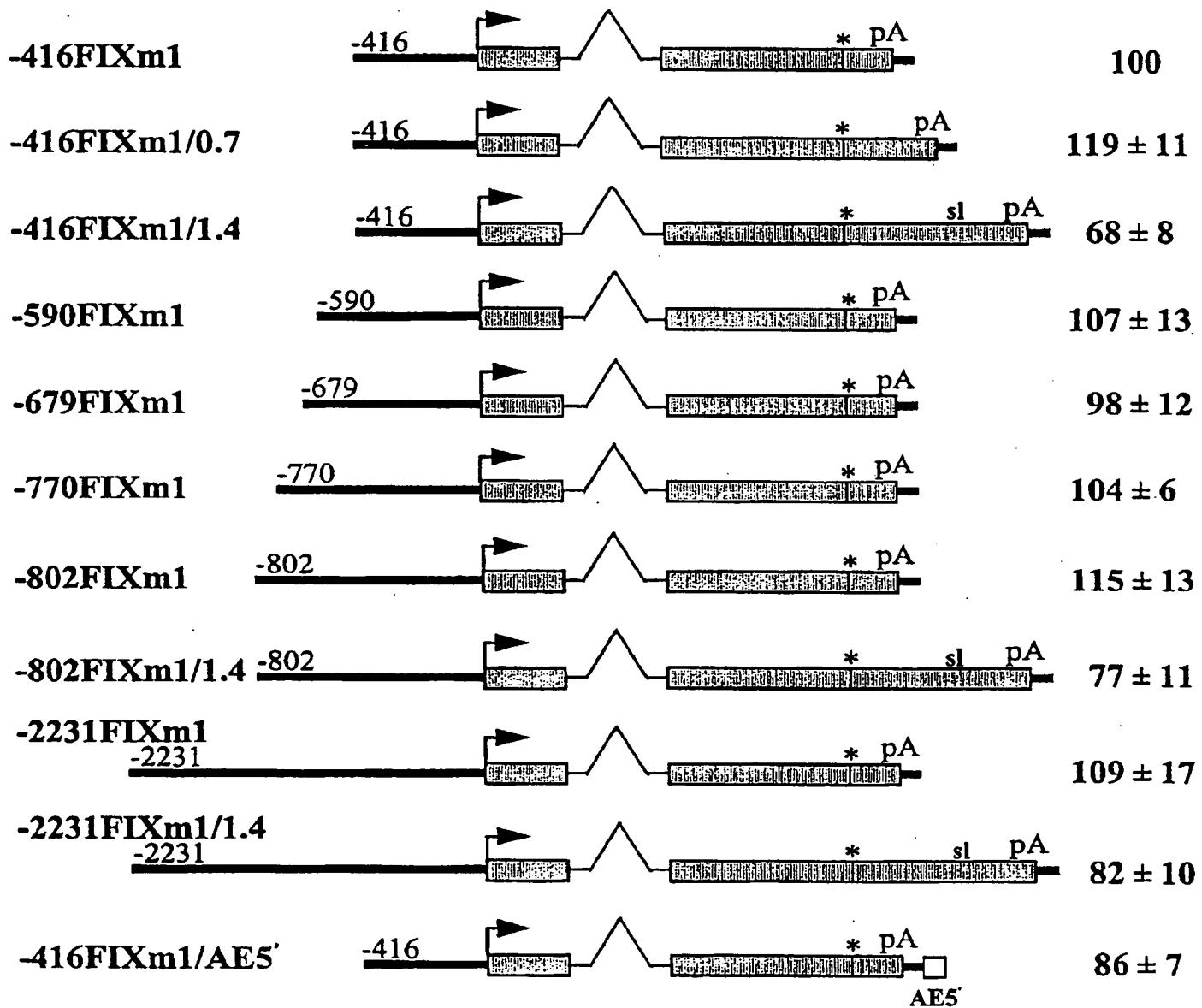
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Figure 1



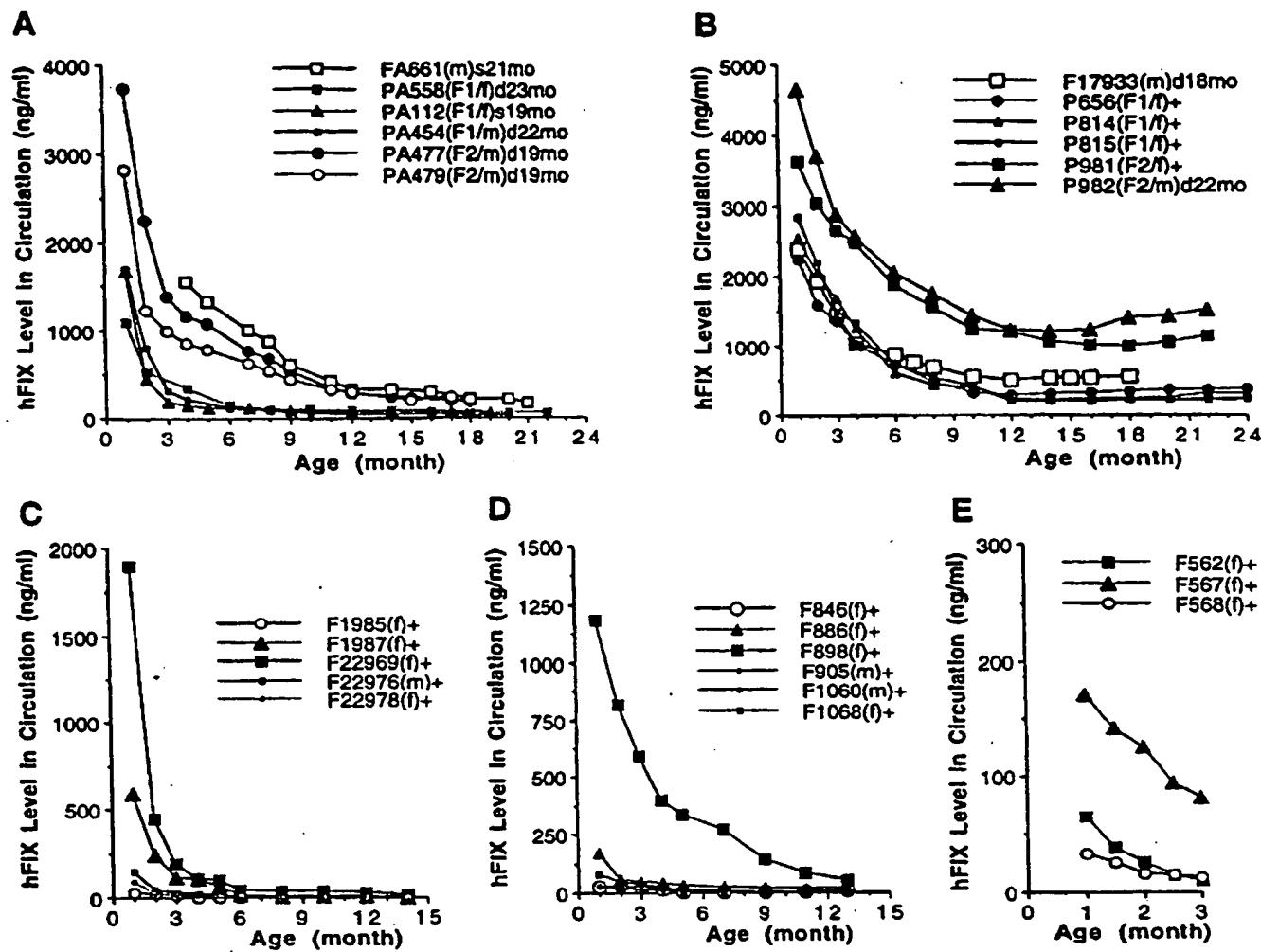
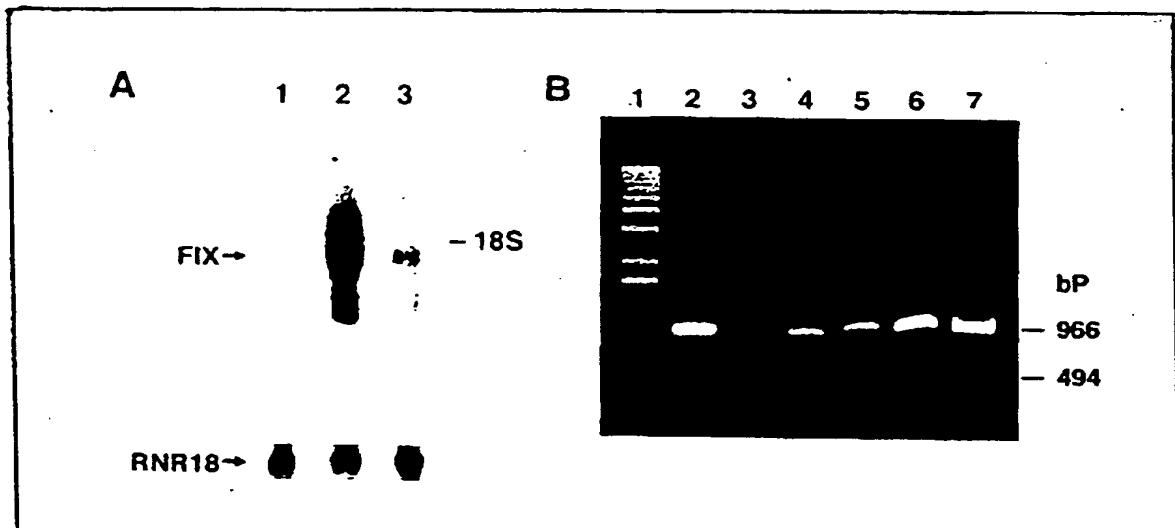


Figure 2

Figure 3

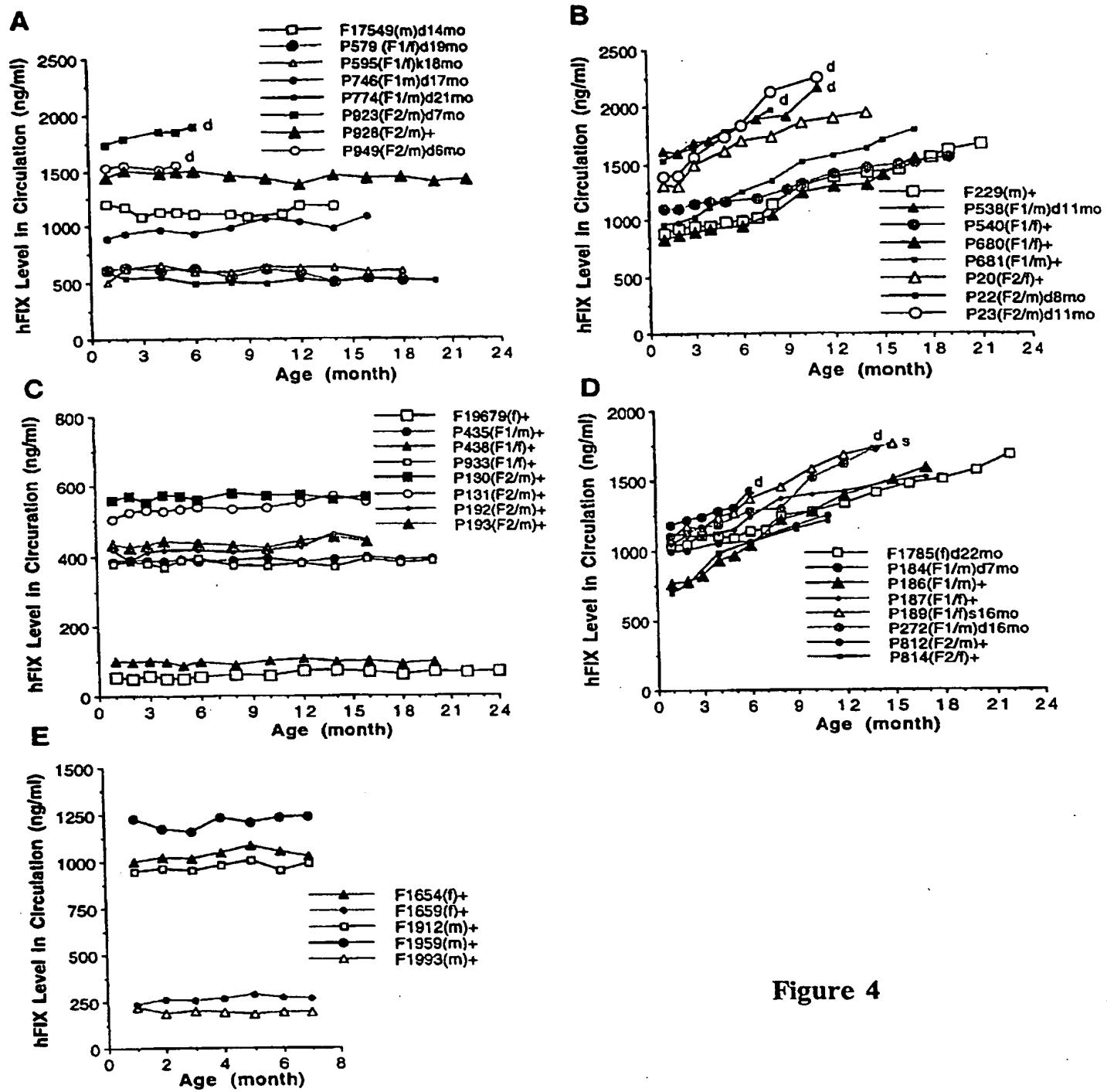


Figure 4

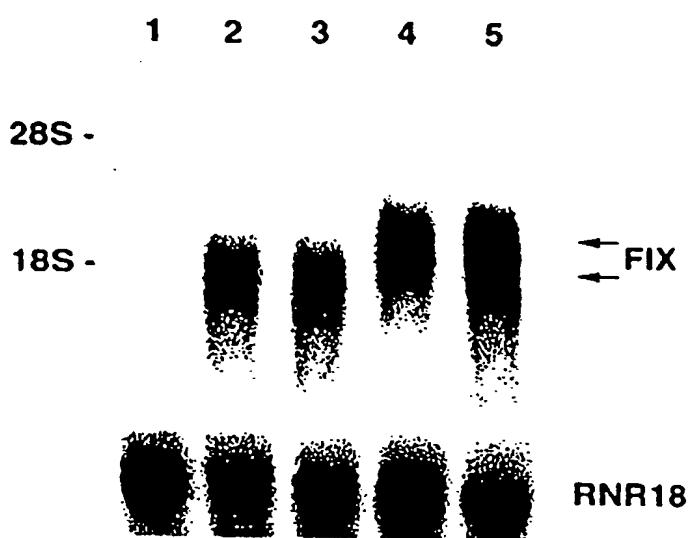
Figure 5

Figure 6

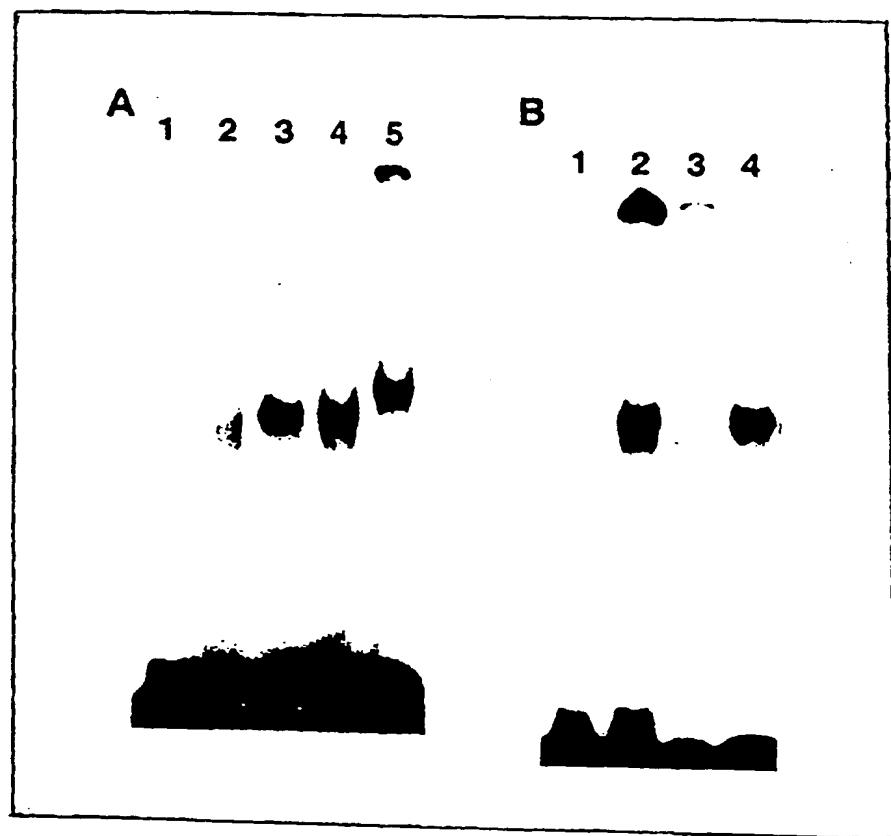


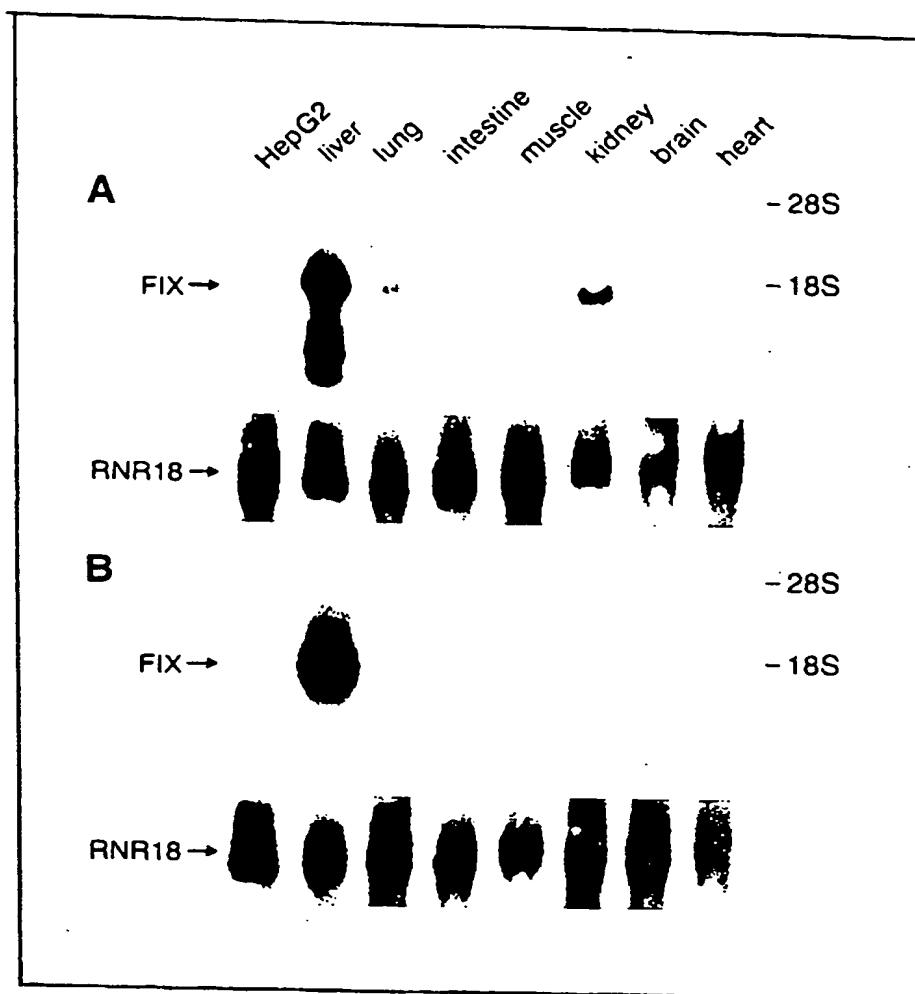
Figure 7

Figure 8A

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CATGGTAGG	AAGAACATCAAT	ATCATGAAA	TGGCCATAAT	GCCAAAGGTA	ATTTATAGAT	TCAATGCCAT	CCCCATCAAG	CTACCAATCA	CTTTCTTCAC	-2566
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-46

Met

32

GATGGACATT	ATTTCCCAAGA	AGTAAATACA	GTCAGCTTG	TACTTTGGTA	CAACTAATCG	ACCTTACAC	TTTCACAAATC	TGCTAGGCAA	GGTT ATG	32
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Gln	Arg	Val	Ash	Met	Ile	Met	Ale	Glu	Ser	Pro	Gly	Leu	Ile	Thi	Ile	Cys	Leu	Leu	Gly	Tyr	Leu	Leu	Ser	Ale	Glu	Cys
CAG	CGC	GTG	AAC	ATC	ATG	GCA	GAA	TCA	CCA	GGC	CTC	ATC	ACC	ATC	TGC	CTT	TTA	GCA	TAT	CTA	CTC	AGT	GCT	GAA	TGT	
-18																										

Thi	ACA	G	TTT	TTTCCCTTTT	TTAAATACAA	TTGAGTATGC	TTGCTCTTAA	GATATAGAAA	TATCTGATGC	TGCTCTCTTC	ACTAAATTTT	GATTACATGA	211
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Figure 8B

Figure 8C

Figure 8D

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 TAATCAGCCTC TGGCTTACAC AAGCTGTTCTG GGAGCACACAT TAACTATCTC TCACTCTCTC TCACTCTCTC TCACTCTCTC 259
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 TGTGTTCTG TGTGAGGAG CCTTCTCTC TGTGAGTGTG TGTGAGAAG GAAATCTCTC 400

Figure 8E

GCTAAATAAA ATTGTTGTTG AATAAATTGG GCTAAAGGCA GAAGGCTCAT AATTTGAGAA CCCACGTCGC ACCGTCCTCC AAGCATCCAT AGTTCTTTG 30329
 ATATACCCCT ATTATCACTT ATTTCAGTGA GGTACAAATTA GTCTCTGATG TAGCCATTTC CATAACAGAA GGCCTTCCCA AAAATCAGTG TCATGTCAAC 30429
 GATCCTTTA TCTCTGGTGC TTGGCACAAAC CTGTACAGG TCCTCAGAAA ACAAAACATTT GAAATTAATGG CCAAATGAGT TTGTCCTCAA AAAAGGGTGC 30529
 AGGATACTTG AAATTTGGAA AATCTAGGAT AATTCATGAC TAGTGGATTC ATTATCACCA ATGAAAGGCT TATAACAGCA TGAGTGAACCA GAACCATCAGC 30629
 TATGATAGTC CTGAAATGGCT TTTTGCTCTG AAAATATGC ATTGGCTCTC ATTACATTTA ACCAAAATTA TCACAAATATA AGAATCAGAT CTITAACATT 30729
 234 Gly Glu
 GCGAAATTAGG TCACTGGTCC CAAAGTAGTCA CTTAGAAAAT CTGTGTATGT GAAATACTGT TTGTCGACTTA AAATGAAATT TATTTTAAT AG GT GAA 30826
 His Asn Ile Glu Glu Thr Glu His Thr Glu Gln Lys Arg Asn Val Ile Arg Ile Ile Pro His His Asn Tyr Asn Ala Ala
 CAT AAT ATT GAG GAG ACA GAA CAT ACA GAG CAA AAC CGA AAT GTG ATT CGA ATT CCT CAC CAC AAC TAC AAT GCA GCT 30907
 Ile Asn Lys Tyr Asn His Asp Ile Ala Leu Leu Glu Leu Asp Glu Pro Leu Val Leu Asn Ser Tyr Val Thr Pro Ile Cys
 ATT AAT AAG TAC AAC CAT GAC ATT GCC CTT CTG GAA CTG GAC GAA CCC TTA GTG CTA AAC AGC TAC GTT ACA CCT ATT TGC 30988
 Ile Ala Asp Lys Glu Tyr Thr Asn Ile Phe Leu Lys Phe Gly Ser Gly Tyr Val Ser Gly Trp Gly Arg Val Phe His Lys
 ATT GCT GAC AAC GAA TAC ACC AAC ATC TTC CTC AAA TTT GGA TCT GGC TAT GTA ACT GCC TGG GGA AGA GTC TTC CAC AAA 31069
 Gly Arg Ser Ala Leu Val Leu Gln Tyr Leu Arg Val Pro Leu Val Asp Arg Ala Thr Cys Leu Arg Ser Thr Lys Phe Thr
 GGG AGA TCA GCT TTT GTT CTT CAG TAC CTT AGA GTT CCA CTT GTT GAC CGA GCC ACA TGT CTT CGA TCT ACA AAG TTC ACC 31150
 Ile Tyr Asn Asn Met Phe Cys Ala Gly Phe His Glu Gly Gly Arg Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro His Val
 ATC TAT AAC AAC ATG TTC TGT GCT GGC TTC CAT GAA GGA GGT AGA GAT TCA TGT CAA GGA GAT AGT GGG GGA CCC CAT GTT 31231
 Thr Glu Val Glu Gly Thr Ser Phe Leu Thr Gly Ile Ile Ser Trp Gly Glu Glu Cys Ala Met Lys Gly Lys Tyr Gly Ile
 ATC GAA GTG GAA GGG ACC AGT TTC TTA ACT GGA ATT ATT AGC TGG GGT GAA GAG TGT GCA ATG AAA GGC AAA TAT GGA ATA 31312
 Tyr Thr Lys Val Ser Arg Tyr Val Asn Trp Ile Lys Glu Lys Thr Lys Leu Thr STOP 415
 TAT ACC AAC GTA TCC CCG TAT GTC AAC TGG ATT AAG GAA AAA ACA AAG CTC ACT TAA TGAAGATGG ATTTCGAGGG TTAATTCACTT 31399
 GGAATTGAAA ATTAACAGGG CCTCTCACTA ACTAAATCACT TTCCCATCTT TTGTTAGATT TGAATATATA CATTCTATGA TCATTGCTTT ITCTCTTAC 31498
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 GAGTGGAGG AAAGTGTGCT AATATAATGC AGTCATTTA ACTTGCTTT TAAGTGTGAT TTGTTTTAGT TTGTTTGAAT ATTAATTTGT TTATACTGAC 34599
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 AACCTGCTC GAATTCTTC TAAACATCTC CTAAATGACCA AAAAGGTGAC GAAATCAAGG AGCAAAATAA GTAGGCCCTG GAAAGCAAGA GTGCC 35094

Figure 9A

Figure 9B

MTKSSHNCLLRPENKPGLWGPQAQAAASLRPSPATLVVSSPGHA
EHPPAAPAQTPGPQVSASARGPGPVAGGSGRMERRMKGGYLDQRVPYTFCSKSPGNGS
LGEALMVPQGKLMDPGSLPPSDSEDLFQDLSHFQETWLAEAQVPDSDEQFVPDFHSEN
LAFHSPTTRIKKEPQSRTDPALSCSRKPPLPYHHGEQCLYSRQIAIKSPAPGAPGQS
PLQPFSSRAEQQSLLRASSSSQSHPGHGYLGEHSSVFQQPVDMCHSFTSPQGGGREPL
PAPYQHQLSEPCPPYPQQNFKQEYHDPLYEQAGQPASSQGGVSGHRYPGAGVVIKQER
TDFAYDSDVPGCASYMLHPEGFSGPSPGDGVMGYGYEKSRLPFPDDVCIVPKKFEGDI
KQEGIGAFREGPPYQRRGALQLWQFLVALLDPTNAHFIAWTGRGMFKLIEPEEVAR
LWGIQKNRPAMNYDKLSRSLRYYEKGIMQKVAGERYVYKFVCEPEALFSLAFPDNQR
PALKAEFDRPVSEEDTVPLSHLDSPAYLPELTGPAPPFGHRRGGYSY

Figure 10 (A)

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 121 ctgctgctca ctcagttcca caggtggag gaacagcagg gcttagatgt ggggtcattg
 181 tgcagatggg aaaacaaagg cccagagagg ggaagaaatg cctaggagct accgagggca
 241 ggcgacccca accacagccc agtgcgtggag ctgtgagatgg atgtagagca
 301 cattcagcca gctcaggggg aggacagggg ccctgaagcc agggatgg
 361 agggagctca gagagaaggg gaggggagtc tgagctcaatg tttccgctgc
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 481 caagtgtggc ccaaccaccc ccaatgtac cagggccatc caccatcccc
 541 atcaggaggg gcttctggg cacaccatc acccgttctt gagtttctt
 601 cattttatc ctcacagcag ctcaacaagg tacataccgt caccatcccc
 661 tagggaaattt gaggctcgga gcggttaaac aactcacccg aggctcaca
 721 gggttccctg gtctgaatgt gtgtgctggg ggatectgtg ggtcactcgc
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 2881 gtcactgttttccatc ttttgcgttgcgtt tggccatgttgcgtt
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Figure 10 (B)

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Figure 10 (C)

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Figure 10 (D)

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Figure 11

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1441 ctggttttta taagagaagt tcctctgaac actattatct tcatgggcag agtagccaac
1501 cttgtgttta agaaaaatgt tcttattttt tgcacctt cctattttg gtttgtgaac
1561 agaagaaaaa ataaatacaa actacttcca tctcacatt

Figure 12 A

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1 ctgcaggggg gggggggggg gggggctgtc atggcggcag gacggcgaac ttgcagtatc
61 tccacgaccc gcccctacag gtgccagtgc ctccagaatg tggcagctca caagcctcct
121 gctgttcgtg gccacctggg gaatttccgg cacaccaggt ccttctgact cagtgttctc
181 cagcagcggag cgtgcccacc aggtgctgcg gatccgcaaa cgtgccaact ctttccttgg
241 ggagctccgt cacagcagcc tggagcggga gtgcatacgag gagatctgtg acttcgagga
301 ggccaaggaa attttccaaa atgtggatga cacactggcc ttctggtcca agcacgtcga
361 cggtgaccag tgcttggct tgcccttggga gcacccgtgc gccagcctgt gctgcggca
421 cggcacgtgc atcgacggca tcggcagctt cagctgcgac tgccgcagcg gctgggaggg
481 cccgcttcgtc cagcgcggagg tgagcttccct caatttgcgtc ctggacaacg gccgcgtcgc
541 gcattactgc ctagaggagg tgggctggcg gcgctgttagc tgtgcgcctg gctacaagct
601 gggggacgac ctcctgcagt gtcaccccgc agtgaagtgc ctttgcgtggga ggcctggaa
661 cgggatggag aagaagcgca gtcacctgaa acgagacaca gaagaccaag aagaccaagt
721 agatccgcgg ctcattgtat ggaagatgac caggcggggaa gacagccccct ggcagggtgt
781 cctgctggac tcaaagaaga agctggcctg cggggcagtg ctcatccacc ctttccttgg
841 gctgacagcg gcccactgca tggatgagtc caagaagctc cttgtcaggc ttggagagta
901 tgacctgcgg cgctgggaga agtgggagct ggacctggac atcaaggagg tcttcgttcca
961 ccccaactac agcaagagca ccaccgacaa tgacatcgca ctgctgcacc tggcccgagcc
1021 cggccaccctc tcgcagacca tagtgcctt catgcctcccg gacagcgcc ttgcagagcg
1081 cgagctcaat caggccggcc aggagacccct cgtgacggc tggggctacc acagcagccg
1141 agagaaggag gccaagagaa accgcacccct cgtcctcaac ttcatcaaga ttccctgtgt
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1381 ctttcacaaac tacggcggtt acacccaaatg cagccgctac ctgcactggc tccatggca
1441 catcagagac aaggaagccc cccagaagag ctgggcaccc ttagcgaccct ccctgcagg
1501 ctgggctttt gcatggcaat ggatgggaca ttaaaggggac atgttacaag cacaccggcc
1561 tgctgttctg tccttccatc cctttttgg gctttctgg agggaaagttaa catttactga
1621 gcacctgttg tatgtcacat gccttatgaa tagaatctt aactccttagag caactctgtg
1681 gggtggggag gagcagatcc aagtttgcg gggtctaaag ctgtgtgtgt tgagggggat
1741 actctgttta taaaaaaagaa taaaaaaacac aaccacgaaa aaaaaaaaaa aaaaaaaaaa
1801 aaaaaaaaaa aaaaaaaaccc ccccccggccc cccccccctg cag

```

Figure 12 B

1 agtgaatctg ggccgagtaac acaaaaacttg agtgtcctta cctgaaaaat agaggtaga
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 121 caattggagg tgagggtgg gcccagtgc cagcacctat gcactgggga cccaaaaagg
 181 agcatcttct catgattta tgtatcagaa attggatgg catgtcattt ggcacagcgtc
 241 tttttcttg tatggtgca cataataca tgtgtctt aattaatgtt attttagatt
 301 tgacgaaata tggaatatta cctgttgtgc tgatcttggg caaactataa tatctctggg
 361 caaaaatgtc cccatctgaa aaacaggag aacgttcctc cctcagccag ccactatggg
 421 gctaaaatgt aaccatctt gtcaagggtt ttgcctcac ctccctccct gctggatggc
 481 atccttgta ggcagagggt ggcttcggc agaacaagcc gtgctgagct aggaccagga
 541 gtgctagtgc cactgtttgt ctatggagag ggaggcctca gtgctgaggg ccaagcaa
 601 atttgtgtt atgattaaac tcgaactcca ggctgtcatg gggcaggac ggcgaacttg
 661 cagtatctcc acgaccggcc cctgtgagtc cccctccagg caggctatg aggggtgtgg
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 781 aggggtctca acaagcctga gcttggggta aaaggacaca aggcctcca caggccaggc
 841 ctggcagcca cagtctcagg tcccttgc atgcgcctcc ctcttccag gccaagggtc
 901 cccaggccca gggccattcc aacagacagt ttggagccca ggaccctcca ttctccac
 961 cccacttcca ccttggggg tgcggattt gaacaaatct cagaagcggc ctcagaggg
 1021 gtcggcaaga atggagagca gggtccggta ggggtgtcag aggccacgtg gcctatccac
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 1201 tcctggctct gcgagtaatg catggatgt aacatggaga cccaggaccc tgcctcagtc
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 1561 ggactaactc cttaattct cacaacaccc ttaaggcac agatacacca cgttattcca
 1621 tccattttac aaatgaggaa actgaggcat ggagcagttt agcatcttgc ccaacattgc
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 3001 ggatgtgtca agatggggct gcatgtggg tactggcaga agatgtgagag gattnaactt
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 3241 catctcagag caaggcttcg tcctccaaact gccatctgtc tcctggggag gaaaagagca
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 3361 agacccccc aggtctccc agtctgtt cctcagaccc cctcatggc ccagcccttc
 3421 ttagggccctt cccaagggtt agtcccttc cttccaaac cagactcgt gttctccagc
 3481 agcgagcgtg cccaccagggt gctgcggatc cgaaacgtg ccaactcctt cttggaggag
 3541 ctccgtcaca gcaagccttgc gccccggatgc atagaggaga tctgtgactt cgaggaggcc

Figure 12 B (continued)

3601 aaggaaattt tccaaaatgt ggatgacaca gtaaggccac catgggtcca gaggatgagg
3661 ctcagggcg agctggtaac cagcaggggc ctcgaggagc aggtggggac tcaatgctga
3721 ggccttta ggagttgtgg gggtggctga gtggagcgat taggatgctg gccctatgat
3781 gtcggccagg cacatgtac tgcaagaaac agaattcagg aagaagctcc aggaaagagt
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4261 aattgtgcac gcctggggcc cttccaagg catccaggaa tgcttccag tggaggctt
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4381 gaccccccatt ctggacctcc atccccacca ctctttcccc cagtggctc cctggcagac
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4681 gaaaaaacact ggcttaggaa ggaggactggaa gggatggagg ctgggcgcgg ggggtggcg
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6961 aaggccgtgg gaaaggccctg tcattggcag aaaaaatgcc tttctgtttt tttatgcatt
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7081 ttcggccctt cagagcaggg tggggcaggg gagctggtgc ctgtgcaggg
7141 tgcattgactc cctgtggta gctaagagca ccactccttc ctgaagcggg

Figure 12 B (continued)

7201 cctagtcaga gcctctgggtt caccttctgc aggcagggag aggggagtca agtcagttag
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Figure 12 B (continued)

10861 tgtctgagaa catgctgtgt gcgggcatcc tcggggaccg gcaggatgcc tgcgagggcg
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10981 gctgggggtga gggctgtggg ctccttcaca actacggcgt ttacacccaaa gtcagccgct
11041 acctcgactg gatccatggg cacatcagag acaaggaaac ccccccagaag agctgggcac
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11161 acatgttaaca agcacacccgg cctgctgttc tgccttcca tccctttt gggcttttct
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11281 taactccttag agcaactctg tgggggtgggg aggagcagat ccaagtttg cggggctaa
11341 agctgtgtgt gttgaggggg atactctgtt tatgaaaaag aataaaaaaac acaaccacga
11401 agccactaga gcctttcca gggctttggg aagagcctgt gcaagccgg gatgctgaag
11461 gtgaggccttgc accagctttc cagctagccc agctatgagg tagacatgtt tagctcatat
11521 cacagaggag gaaactgagg ggtctgaaag gtttacatgg tggagccagg attcaaatct
11581 aggtctgact cccaaaccca ggtgttttt tctgttctcc actgtcctgg aggacagctg
11641 tttcgacggt gctcgtgtg gaggccacta ttagctctgt agggaaagcag ccagagaccc
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Figure 13 (A)

SEQ ID NO:3

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tagaaaatgg aaccactaga ggaatataat gtgttaggaa attacagtca tttctaaggg
cccagccctt gacaaaatgg tgaagttaaa ttctccactc tgcacatcg atactatgg
tctccactat ggcaactaac tcactcaatt ttccctccctt agcagcattc catctcccg
atcttctttg cttctccaaac caaaacatca atgtttatta gtctgtata cagtaggaa
tctttgtct actctatcac aaggccagta ccacactcat gaagaaagaa cacaggagta
gctgagagggc taaaactcat caaaaacact actcctttc ctctacccta ttccctcaatc
tttacctt tccaaatccc aatccccaaa tcagttttc tctttcttac tccctctctc
cctttaccc tccatggcg ttaaaggaga gatggggagc atcattctgt tataacttctg
tacacagtta tacatgtctc tcaaaccagg acttgcttcc atagtggaga cttgctttc
agaacatagg gatgaagtaa ggtgcctgaa aagttgggg gaaaagttt ttcagagag
ttaagttatt ttatataat aatataatata taaaatataat aatataacaat ataaatataat
agtgtgtgtg tgtatgcgtg tgttagaca cacacgcata cacacatata atggaagcaa
taagccattc taagagctt gatggttatg gaggtctgac taggcatgat ttacacgaa
caagattggc atatcattgt aactaaaaaa gctgacattt acccagacat attgtactct
ttctaaaaat aataataata atgctaacag aaagaagaga accgttcgtg tgcaatctac
agctagtaga gactttgagg aagaattcaa cagtgtgtct tcagcagtgt tcagagccaa
gcaagaagtt gaagttgcct agaccagagg acataagtat catgtctcct ttaactagca
taccccgaa tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
ctaagttgtc cttttctgg ttcgtgttca ccatgaaaca ttttgattat agttaatcct
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SEQ ID NO:76

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tagaaaatgg aaccactaga ggaatataat gtgttaggaa attacagtca tttctaaggg
cccagccctt gacaaaatgg tgaagttaaa ttctccactc tgcacatcg atactatgg
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tctttgtct actctatcac aaggccagta ccacactcat gaagaaagaa cacaggagta
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gcaagaagtt gaagttgcct agaccagagg acataagtat catgtctcct ttaactagca
taccccgaa tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
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tctatcttga atctt

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Figure 13 (B)

SEQ ID NO:77

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tagaaaatgg aaccactaga ggaatataat gtgttaggaa attacagtca tttctaaggg
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atcttcttg cttctccaaac caaaacatca atgttattt gttctgtata cagtaggaa
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cctttaccc tccatggcg ttaaaggaga gatggggacg atcattctgt tataactctg
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agtgtgtgtg tgtatgcgtg ttttttttgcata cacacgcata cacacatata atggaaagcaa
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gcaagaagtt gaagttgcct agaccagagg acataaagtat catgtctc ttaacttagca
tacccgaag tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
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tctatcttga atctt

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SEQ ID NO:78

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atcttcttg cttctccaaac caaaacatca atgttattt gttctgtata cagtaggaa
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gctgagaggg taaaactcat caaaaacact actcctttc ctctacccta ttcctcaatc
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gcaagaagtt gaagttgcct agaccagagg acataaagtat catgtctc ttaacttagca
tacccgaag tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
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Figure 13 (C)

SEQ ID NO:79

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cccagccctt gacaaaattg tgaagttaaa ttctccactc tgccatcg atactatgg
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tacccgaag tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
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SEQ ID NO:80

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atttcttttg cttctccaac caaaacatca atgtttatta gttctgtata cagtacagga
tctttggct actctatcac aaggccagta ccacactcat gaagaaagaa cacaggagta
gtcgagggc taaaactcat caaaacact acetcctttc ctctacccta ttccctcaatc
tttacctt tccaaatccc aatccccaaa tcagttttc tctttcttac tccctctc
cctttaccc tccatggcg taaaaggaga gatggggagc atcattctgt tatacttctg
tacacagtta tacatgtcta tcaaaccagg acttgcttc atagtggaga cttgcttcc
agaacatagg gatgaagtaa ggtgcctgaa aagtttgggg gaaaagttc tttcagagag
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taagccattc taagagctt gatggttatg gaggtctgac taggcgtat ttacacgaaagg
caagattggc atatcattgt aactaaaaaa gctgacattt acccagacat attgtactct
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gcaagaagtt gaagttgcct agaccagagg acataagtat catgtctcct ttaactagca
tacccgaag tggagaaggg tgcagcaggc tcaaaggcat aagtcttcc aatcagccaa
ctaagttgtc cttttcttgc ttcgtgttca ccatggaaaca ttttgattat agttatcct
tctatcttga atctt

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Figure 13 (D)

SEQ ID NO:81

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tagaaaatgg aaccactaga ggaatataat gtgttaggaa attacagtca tttctaaggg
cccagccctt gacaaaattg tgaagttaaa ttctccactc tgtccatcag atactatgg
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SEQ ID NO:82

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Figure 13 (E)

SEQ ID NO:83

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tctatcttga atctt

Figure 14

GAATTCGTAA AGCATTCCCT ATGGTGTACCT GCCCCCTGGGC AAGGTGGGCC TGACTTGTAA -1403
 GAGTGTAGA GTTTTACCCCT GTTCCCTCTAG GAGGGCCTGG TACCAACACA GCCCAGCATG -1343
 GTGTGGTCCC TCAGCAGGAG GCATCTGGTT ACAATCAACA CAAGCTGTC CAGCCAATT -1283
 AAAGAAACCT CAGGAGGAAT AGGGTTTAG GAGGGCATGG GGACCCCTCCT GCACCCGAAG -1223
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 CTTCCCTGTG CAATCAATGG AAGCCACCGA GGCAGGGTCA CATGCCCTGTTAGAGGTGC -923
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 CAAATATTG TGGTTATGGA TTAACCTCGAA CTCCAGGCTG TCATGGGGC AGGACGGCGA +38
ACTTGCAGTC TCTCCACGAC CGGCCCCCTGT GAGTCCCCCT CCAGGCAGGT CTATGAGGGG +98
 TGTGGAGGGA GGGCTGCCCT CGGGAGAAGA -----
 -----1350 bp-----

 ----- MET TRP GLN LEU -39
 ----- AAGAAGTCCT CCTCAGACAG GTGCCAGTGC CTCCAGAATG TGG CAG CTC +1527

Figure 15

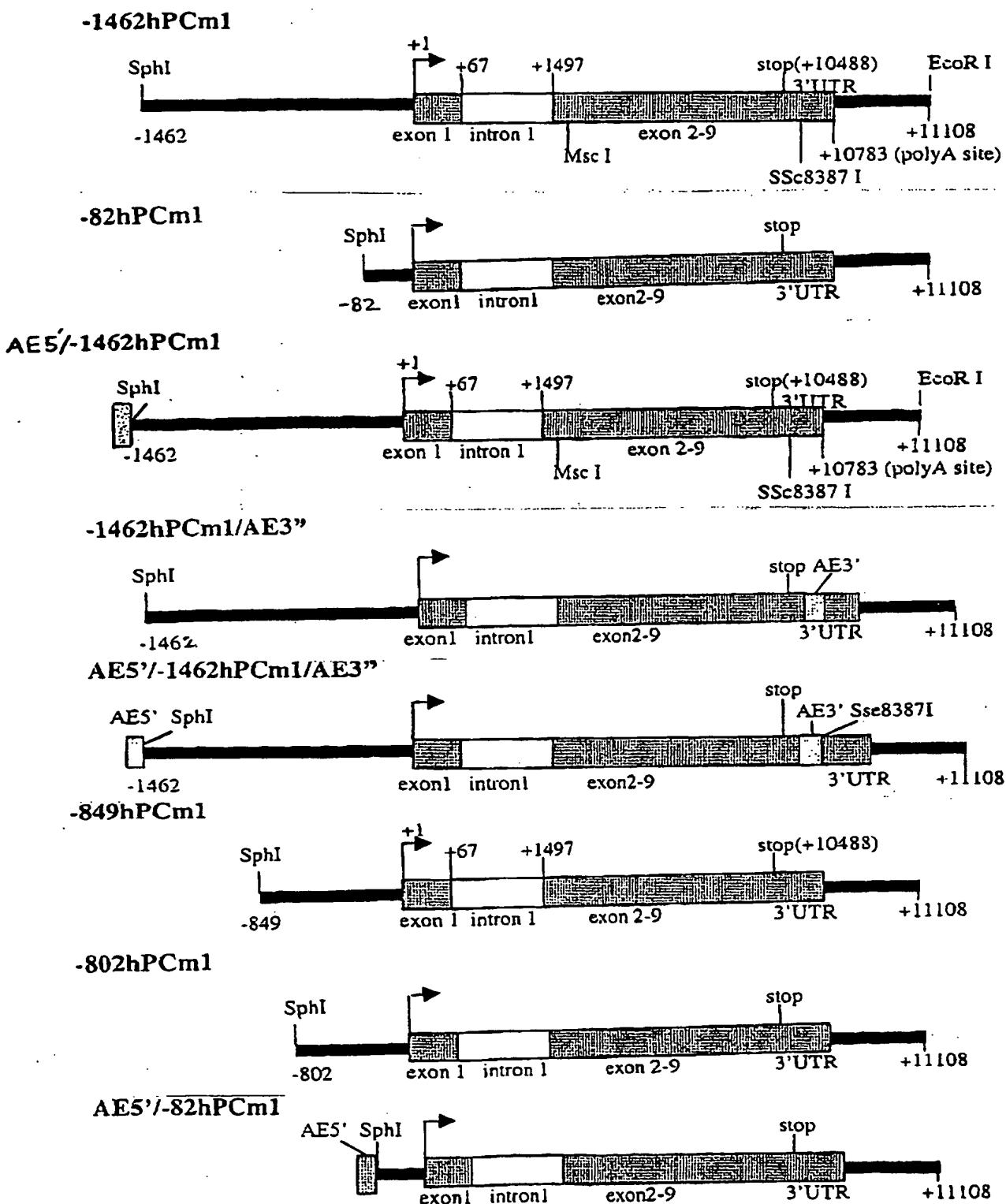


Figure 16

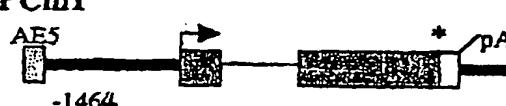
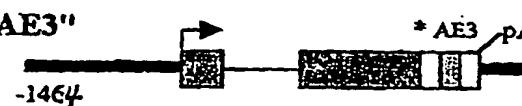
<u>hPC Minigene Constructs</u>	<u>Expression Activity</u> (% \pm SD)
-1464hPCm1 	100
-82hPCm1 	98.7 \pm 11.8
AE5'-1464hPCm1 	101.9 \pm 12.5
-1464hPCm1/AE3'' 	70.1 \pm 7.5
AE5'-1464hPCm1/AE3'' 	74.0 \pm 3.8

Figure 17A

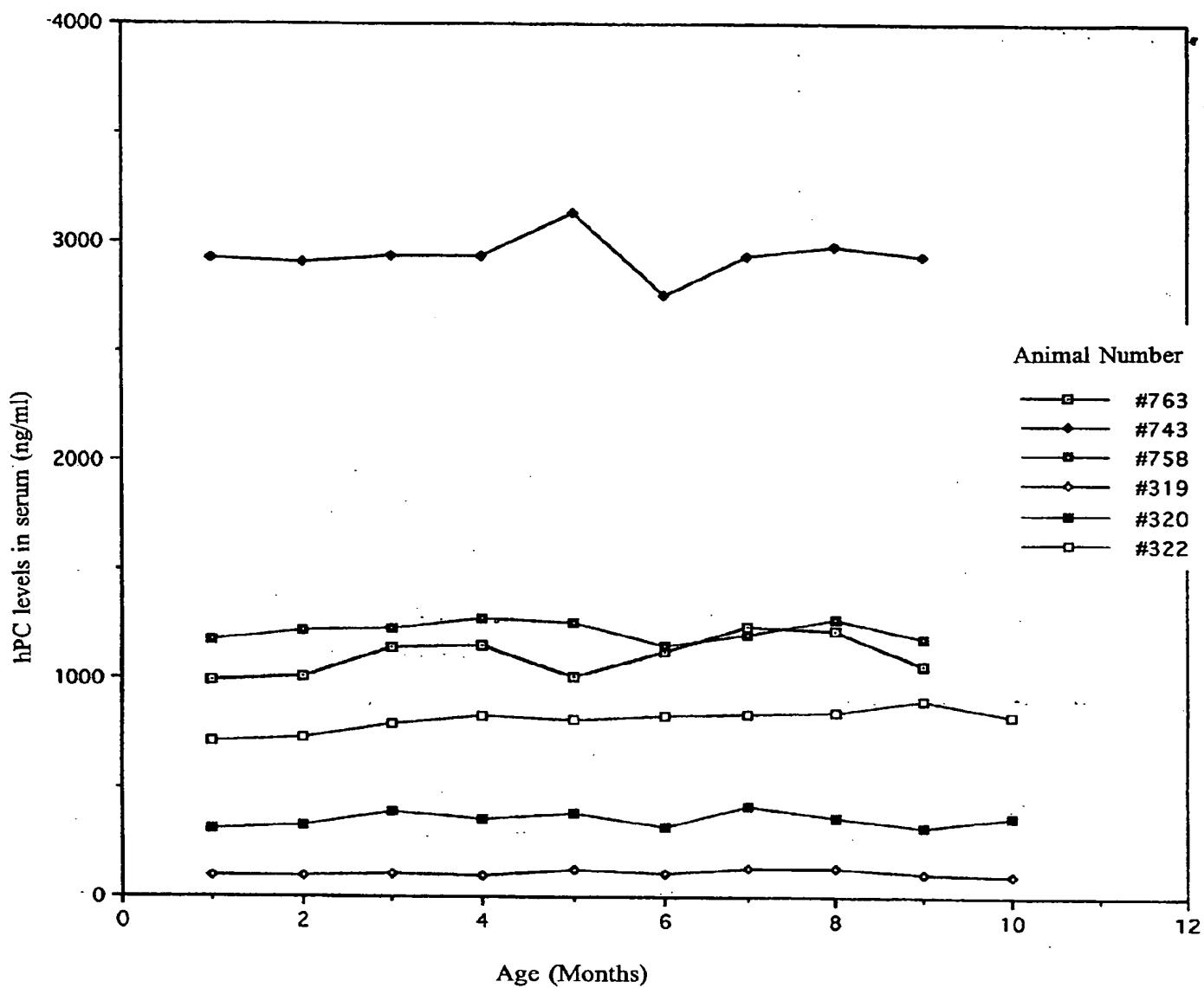


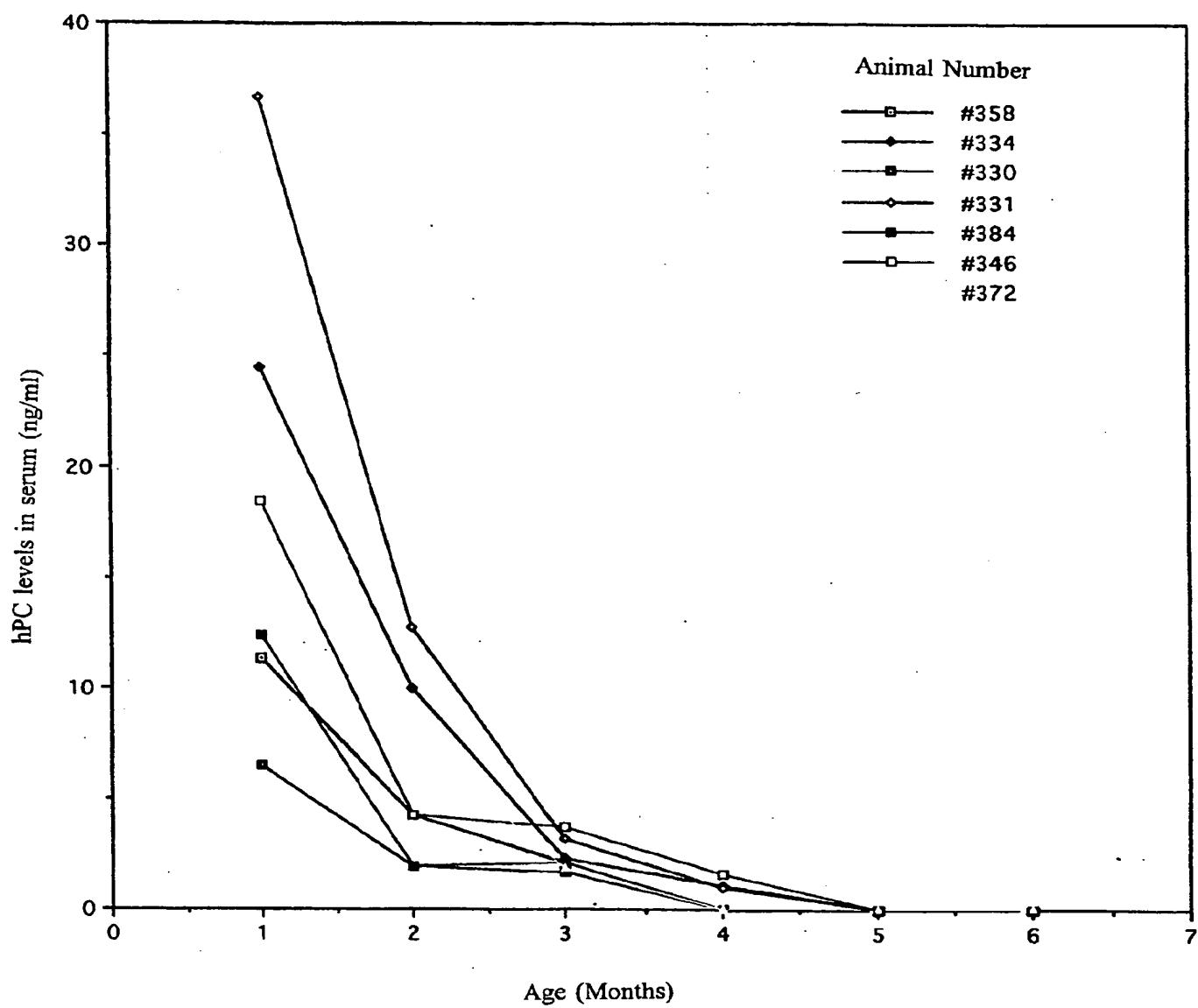
Figure 17B

Figure 17C